

CRF Errors Edited by the STIC Systems Branch

Serial Number: 10/522,991 : CRF Edit Date: 2/1/05
Edited by: NC

_____ Realigned nucleic acid/amino acid numbers/text in cases where the sequence text "wrapped" to the next line

_____ Corrected the SEQ ID NO. Sequence numbers edited were:

_____ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

Deleted: invalid beginning/end-of-file text ; page numbers

_____ Inserted mandatory headings/numeric identifiers, specifically:

_____ Moved responses to same line as heading/numeric identifier, specifically:

_____ Other:



PCT

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/522,991

DATE: 02/11/2005
TIME: 16:11:15

Input Set : A:\PTO.YF.txt
Output Set: N:\CRF4\02112005\J522991.raw

3 <110> APPLICANT: Hitachi High-Technologies Corporation
5 <120> TITLE OF INVENTION: A method of detecting a nucleic acid using DNA
6 microarray and a nucleic acid detecting apparatus
8 <130> FILE REFERENCE: PH-1503-PCT
C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/522,991
C--> 10 <141> CURRENT FILING DATE: 2005-02-02
10 <160> NUMBER OF SEQ ID NOS: 4
12 <170> SOFTWARE: PatentIn Ver. 2.0

Does Not Comply
Corrected Diskette Needed
(PS-1)

ERRORED SEQUENCES

47 <210> SEQ ID NO: 4
48 <211> LENGTH: 50
49 <212> TYPE: DNA
50 <213> ORGANISM: Artificial Sequence
52 <220> FEATURE:
53 <223> OTHER INFORMATION: Description of Artificial Sequence:synthetic DNA
55 <400> SEQUENCE: 4 50
56 tcctctaatt ccaggatcaa caacaaccag aggtttgca tggtcccgta
E--> 62 1/2
E--> 63 dsmdb.1878912.1

Deleted

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/522,991

DATE: 02/11/2005
TIME: 16:11:16

Input Set : A:\PTO.YF.txt
Output Set: N:\CRF4\02112005\J522991.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:62 M:254 E: No. of Bases conflict, LENGTH:Input:2 Counted:51 SEQ:4
L:62 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
M:254 Repeated in SeqNo=4
L:63 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:9
L:63 M:112 C: (48) String data converted to lower case,
L:63 M:252 E: No. of Seq. differs, <211> LENGTH:Input:50 Found:58 SEQ:4
L:63 M:252 E: No. of Seq. differs, <211> LENGTH:Input:50 Found:58 SEQ:4